

Fig. 1a

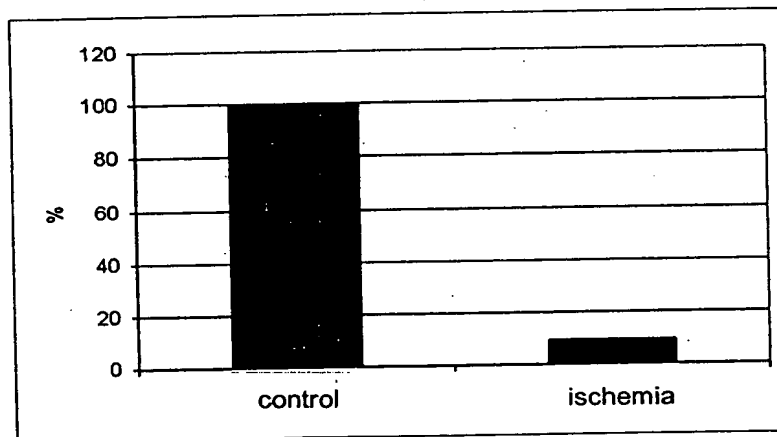


Fig. 1b

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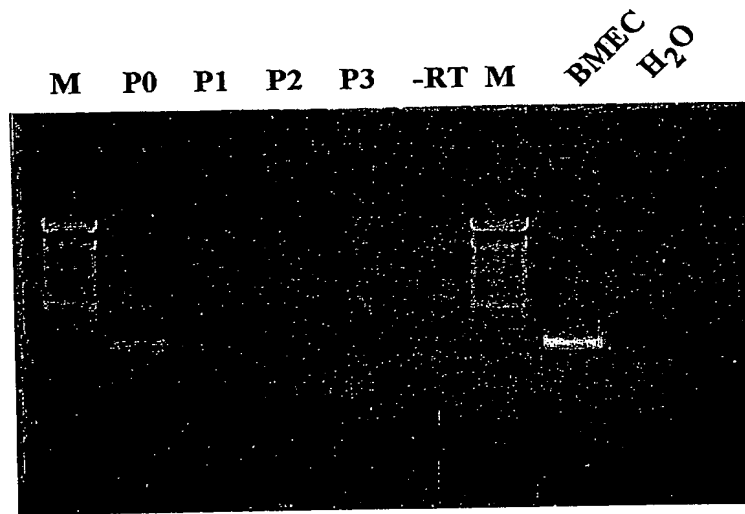


Fig. 2

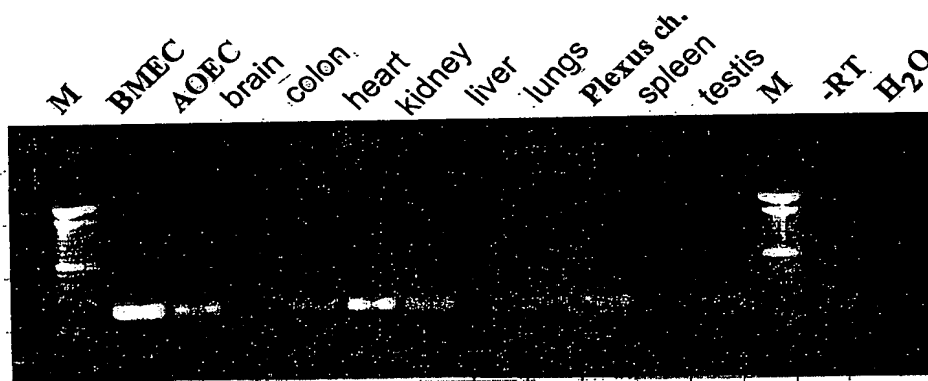


Fig. 3

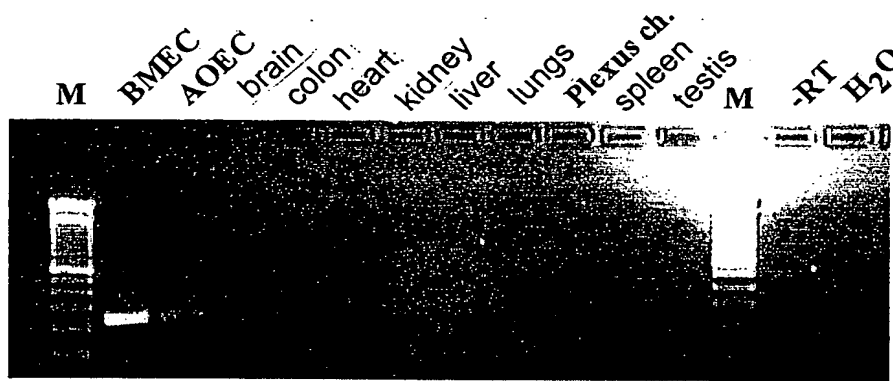


Fig. 4

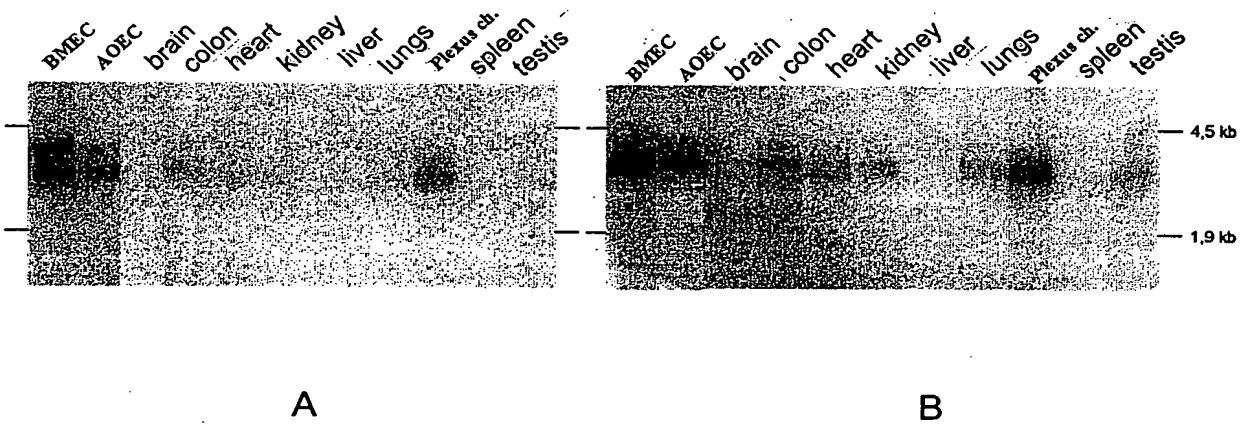


Fig. 5

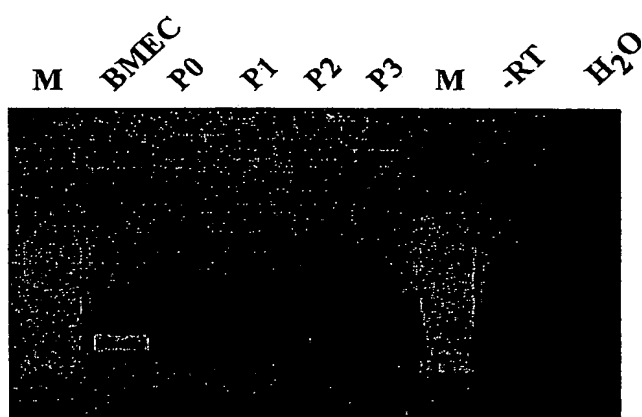


Fig. 8

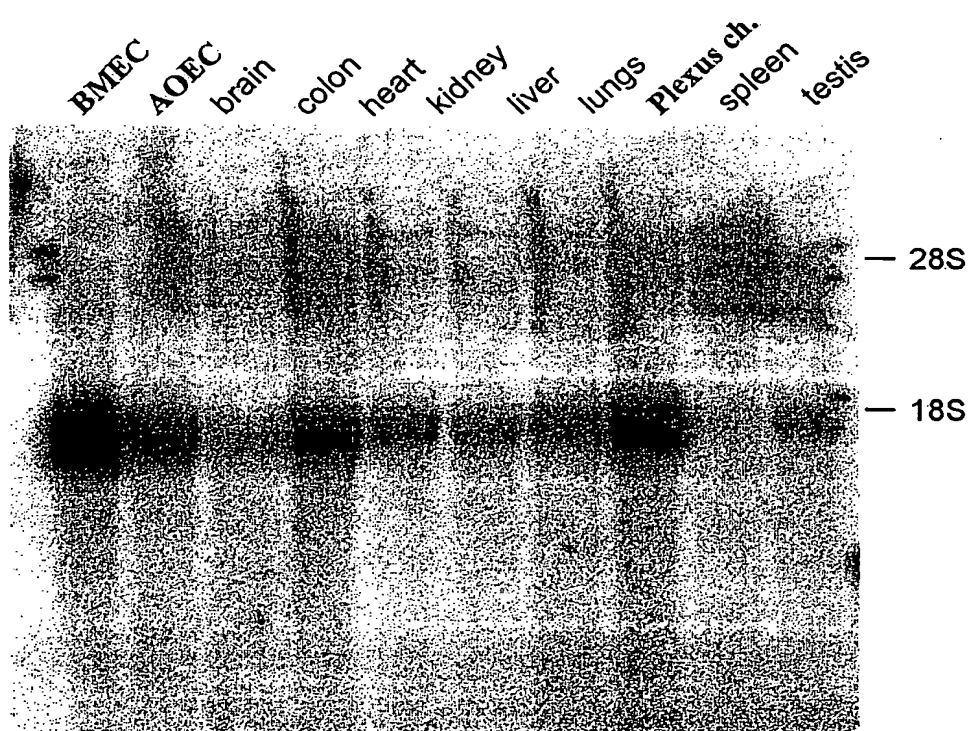


Fig. 9

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human      M--AARTGHTALRRVVSGCRPKSATAAG----AQAPVRNGRYLASCGILMSRTLPLHTSI 54
pig        MFVAARTGQRTLRLKVVSGCRPKSATATGVPAPAQGPPIRNYLASCGILMNRTLPLHSSF 60
mouse      MIMAARTSQRALARVASGCHPKSTTVTEAP--ARGSARDVRHLAACGVLINRTLPPCAAV 58
          *  ****:::* :*.****:***:.*:  *:. . *: :***:***:.*.***** :..

human      LPKEICARTFFKITAPLINKRKEYSERRILGYSMQEMYDVVSGVEDYKHFPWCKKSDVI 114
pig        LPKEMYARTFFRIAAPLINKRKEYSERRIIGYSMQEMYDVVSGMEDYKHFPWCKKSDVI 120
mouse      LPKEICARTFFRISAPLVNKRKEYSERRILGYSMQEMYDVVSGMEDYQHFPWCKKSDII 118
          ****: *****:*.****:*****:*****:*****:***:*****:*.

human      SKRSGYCKTRLEIGFPPVLERYTSVVTLVKPHLVKASCTDGRLFNHLETIWRFPGLPGY 174
pig        SRRSGYCKTRLEIGFPPVLERYTSVVTLVKPHLVKASCADGKLFNHLETVWRFPGLPGY 180
mouse      SRRSGYCKTRLEVGFPPVLERYTSIVTLVKPHLVKASCTDGKLFNHLETIWRFPGLPGY 178
          *:*****:*****:*****:*****:*****:***:*****:*****

human      PRTCTLDFSISFEFRSLLHSQATLFFDEVVKQMVAAFERRACKLYGPETNIPRELMLHE 234
pig        PRTCTLDFSISFEFRSLLHSQATLFFDEVVKQMVAAFERRACKLYGPETSIPRELMLHE 240
mouse      PRTCTLDFSISFEFRSLLHSQATLFFDEVVKQMVAAFERRACKLYGPETNIPRELMLHE 238
          *****:*****:*****:*****:*****:*****:*****

human      VHHT 238
pig        VHHT 244
mouse      IHHT 242
          :***

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Fig. 10

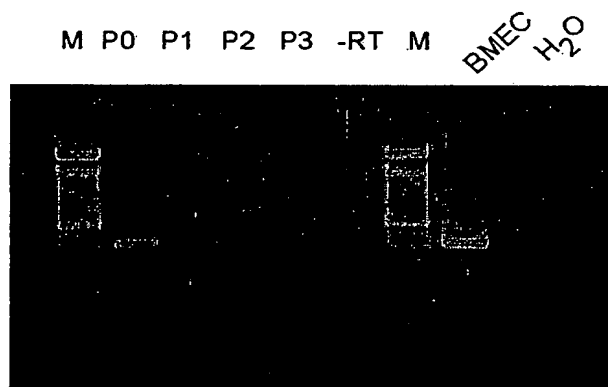


Fig. 11

MGNQVEKLTH	LSYKE <u>EVPTAD</u>	<u>PTGVDRDDGP</u>	<u>RIGVSYIFSN</u>	DDEDVEPQPP
PQGPDGGGLP	DGGDGPPPPQ	PQPYDPR <u>LHE</u>	<u>VECSVFYRDE</u>	<u>CIYQKSFAPG</u>
<u>SAALSTYTPE</u>	<u>NLLNKCKPGD</u>	LVEFVSQAQY	PHWAVYVGNF	QVVHLHRLEV
INSFLTDASQ	GRRGR <u>VVNDL</u>	<u>YRYKPLSSSA</u>	VVRNALAHVG	AK <u>ERELSWRN</u>
<u>SESFAAWCRY</u>	GKREFKIGGE	LR <u>IGKQPYRL</u>	<u>QIQLSAQRSH</u>	<u>TLEFQSLEDL</u>
<u>IMEKRRNDQI</u>	GRAAVLQELA	THLHPAEPEE	GDSNVARTTP	PPGRPPAPSS
EEEDGEAVAH				

Fig. 12

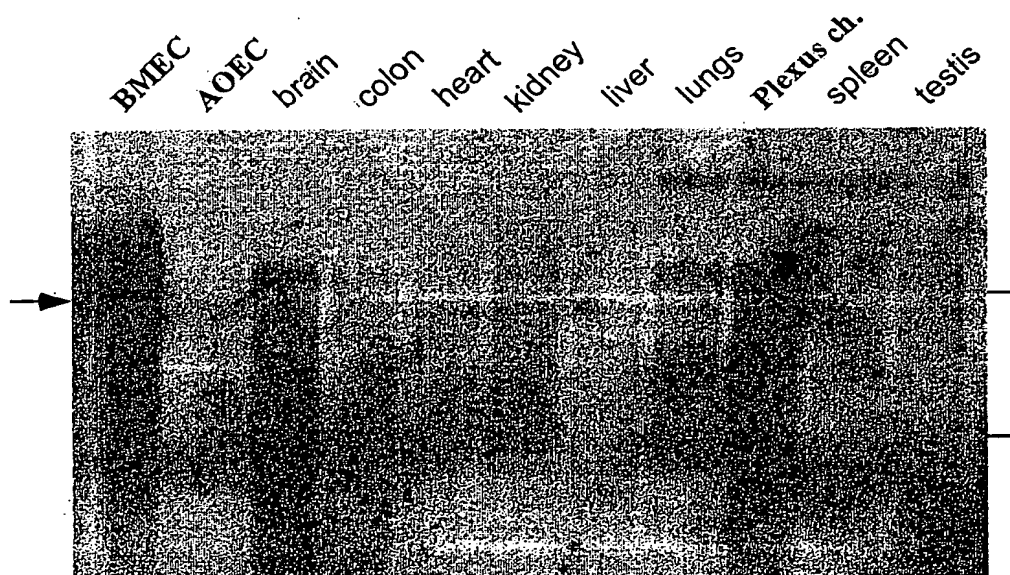


Fig. 13


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MGNQVEKLTHLSYKEVPTADPTGVDRDDGPRIGVSYIFSNDDEDVEPQPP 50
-----
PQGPDGGGLPDGGDGPPPPQPPYDPRLEHEVECSVFYRDECIYQKSFAPG 100
+++++
SAALSTYTPENLLNKCKPGDLVEFVSQAQYPHWAVYVGNFQVVHLHRLEV 150
-----
INSFLTASQGRGRVNDLYRYKPLSSAVVRNALAHVGAKERELSWRN 200

SESFAAWCRYGKREFKIGGELRIGKQPYRLQIQLSAQRSHTLEFQSLEDL 250

IMEKRRNDQIGRAAVLQELATHLHPAEPEEGDSNVARTTPPPGRPPAPSS 300
+++++
EEEDGEAVAH 310
+++++

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possible PEST sequences
poor PEST sequences

Fig. 16

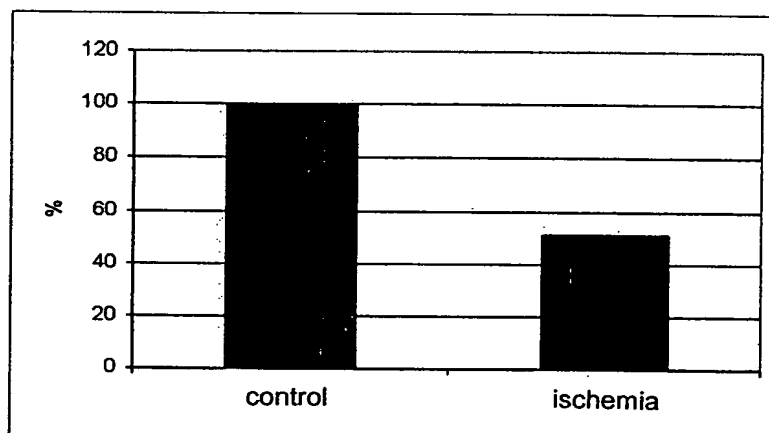


Fig. 17

Fig. 18

Fig. 19

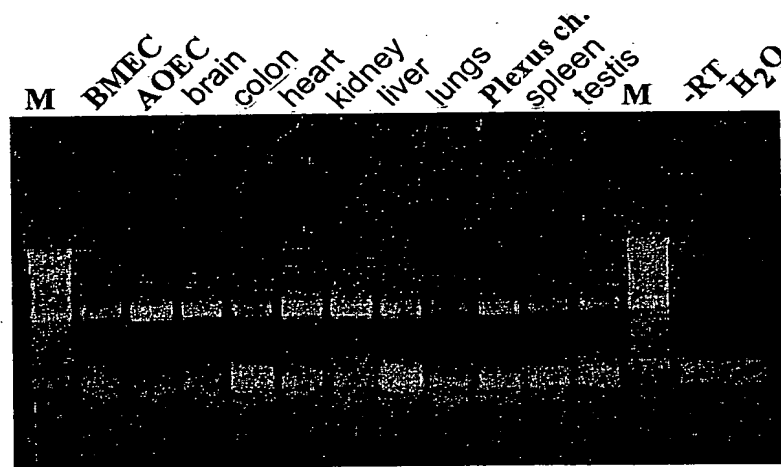


Fig. 20

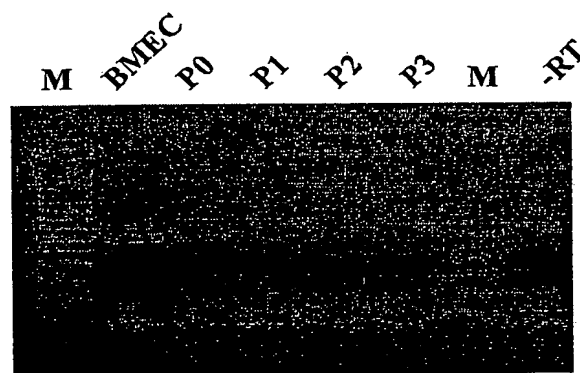


Fig. 21

<u>MAAPEPLRPR</u>	LCRLVRGEQG	YGFHLHGEKG	<u>RRGQFIRRV</u>	PGSPAEEAAL
AGDRLVEVNG	<u>VNVEGETHHQ</u>	<u>VVQRIKAVEG</u>	QTR <u>LLVVDQE</u>	<u>TDEELRRRQL</u>
TCTEEMAQRG	LPPAHDPWEP	KPDWAHTGSH	SSEAGKKDVS	GPLRELRPRL
CHLR <u>KGPQGY</u>	<u>GFNLHSDKSR</u>	<u>PGQYIRSVDP</u>	GSPAARSLR	AQDR <u>LIEVNG</u>
<u>QNVGLRHAE</u>	<u>VVASIKARE</u>	EARLLVVDPE	TDEHFKRLRV	TPTEEHVEGP
LPSPVTNGTS	PAQLNGGSAC	SSRSDLPGSD	KDTE <u>DGSAWK</u>	<u>QDPFQESGLH</u>
<u>LSPTAAEARR</u>	<u>RLEPCESTSA</u>	<u>RHRWTGTGSV</u>	KSSATSEPLP	ACLGTGLGPLP
HGPWASACPE	LPQPQWTGGW	SCHCPEISPS	PGEPPSCPCP	PGTGGLWQQD
RGRETQRCER	ESETETERER	ERHRERQRES	ERARGSRGAR	AFAALPGPAD

Fig. 22

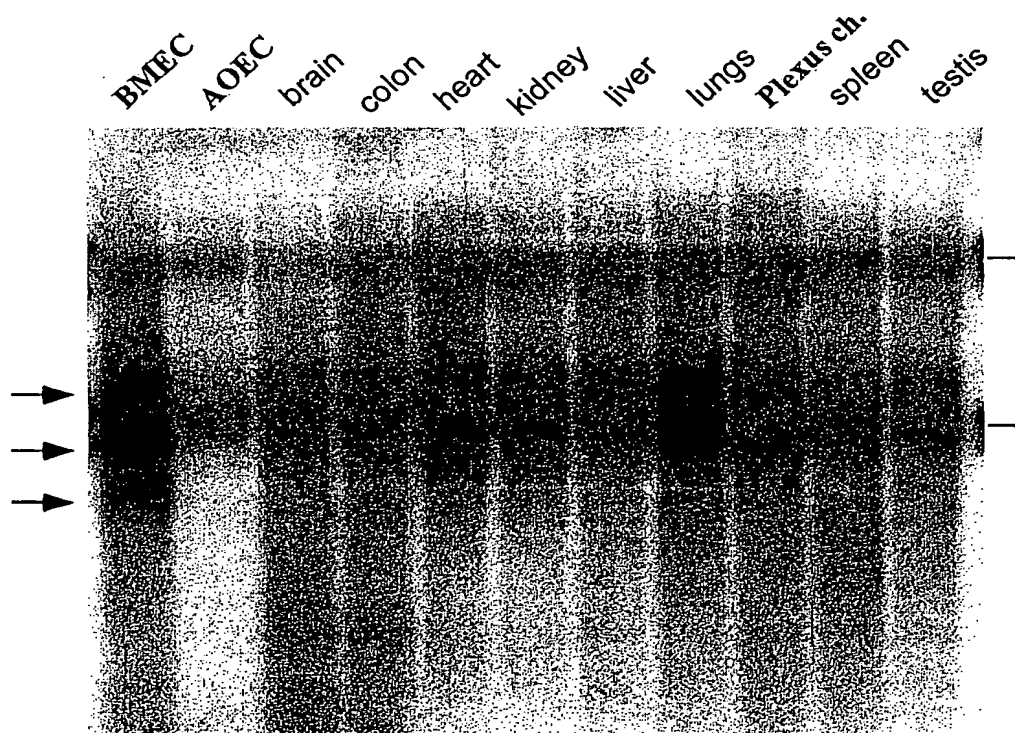


Fig. 23

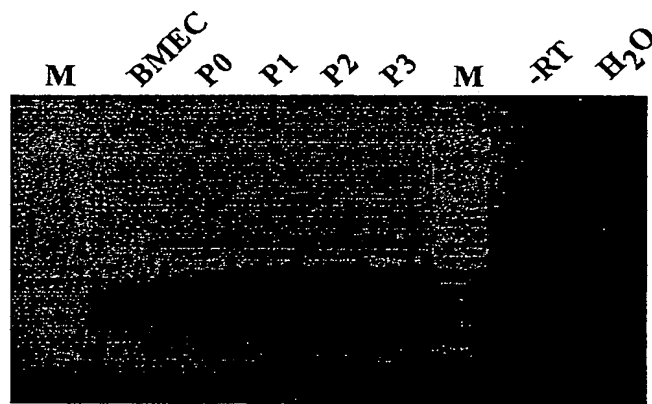


Fig. 24

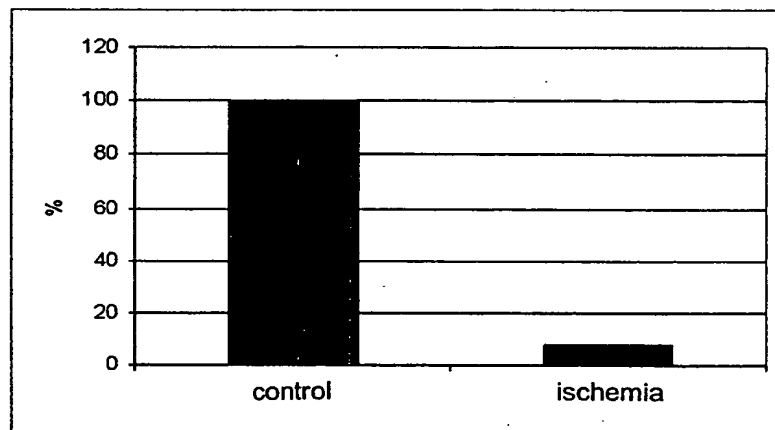


Fig. 25

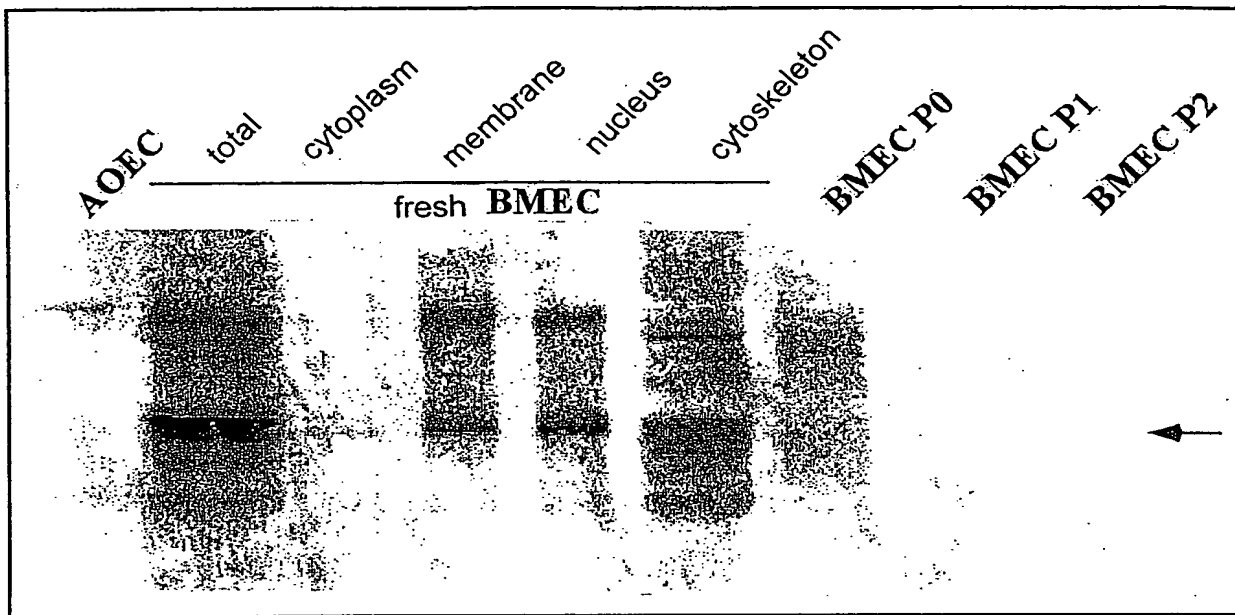


Fig. 26

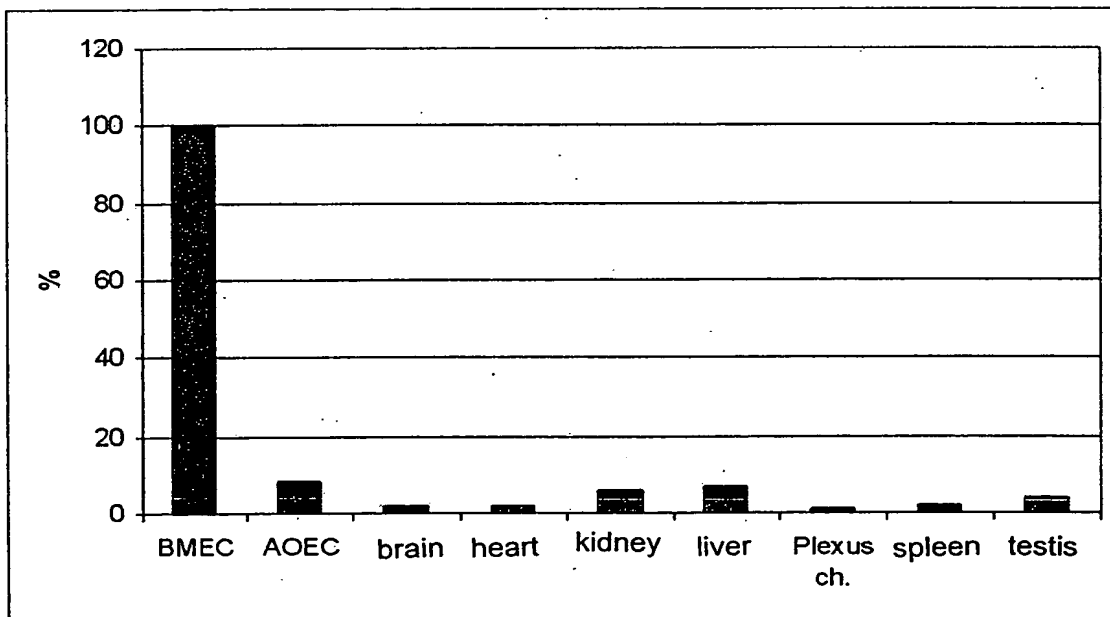


Fig. 27

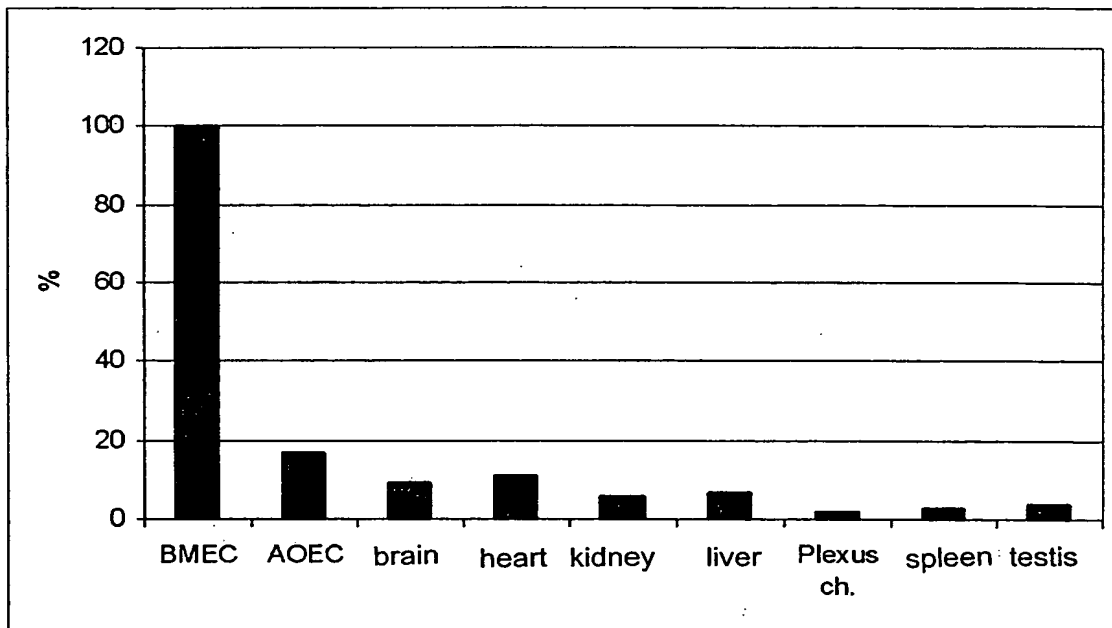


Fig. 28

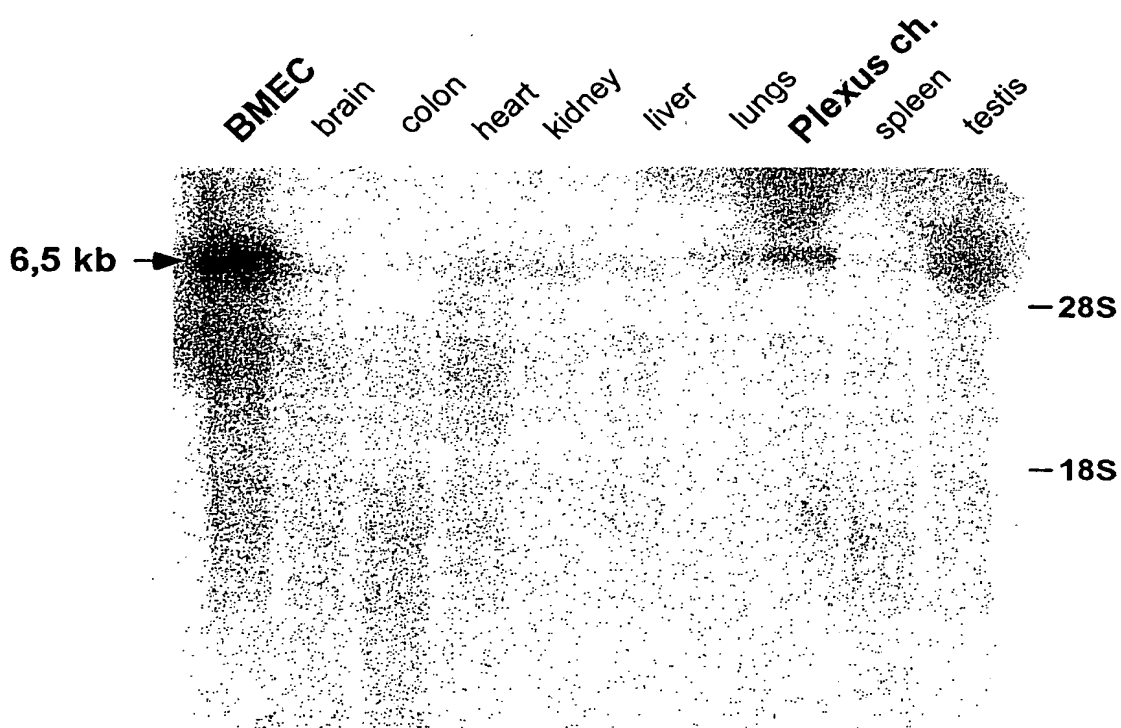


Fig. 29

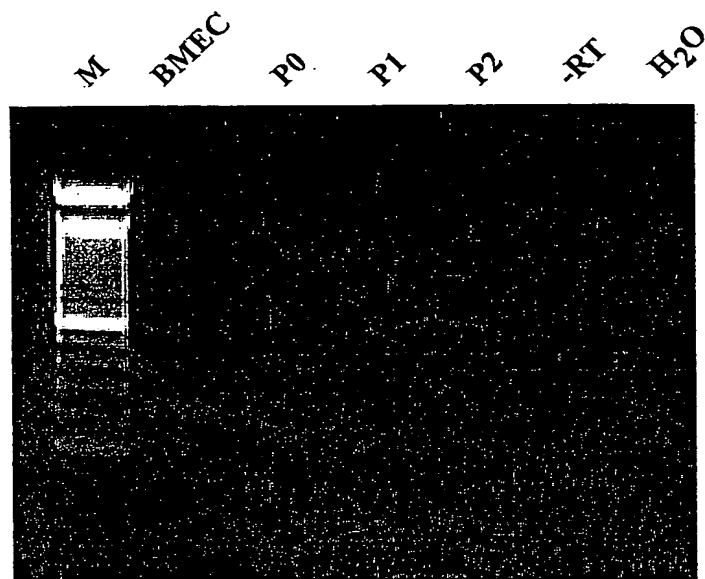


Fig. 30

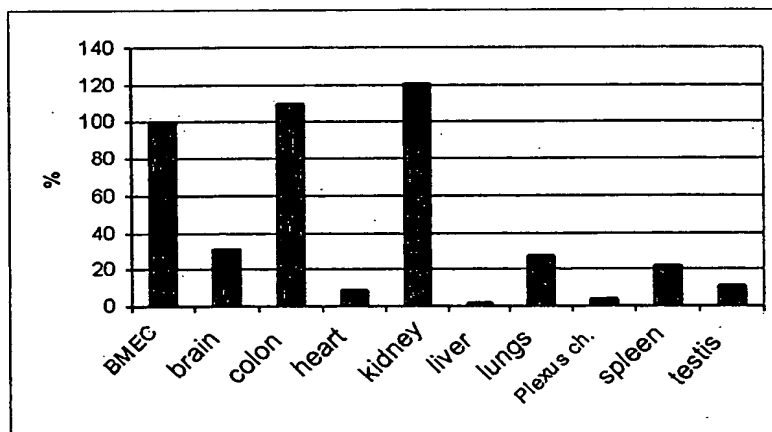


Fig. 31

HSNV1: 48 MSLCICGTAITSQYLAERYKVNTPMLQSFINYCLLFliYTVMLAFRSGSDNLLVILKRK 107
 MSLCICGTAITSQYLAE+YKVNTPMLQSFINYCLLFliYT+MLAF+SG++NLL ILK+K
 PNOV1: 1 MSLCICGTAITSQYLAERYKVNTPMLQSFINYCLLFliYTMMLAFQSGNLLCILKKK 60

 HSNV1: 108 WWKYILLGLADVEANYVIVRAYQYTTLTsvQLLDCFGIPVLMALSWFILHARYRVIHFIA 167
 WWKYILLGLADVEANY+IVRAYQYTTLTsvQLLDCFGIPVLMALSWFIL+ARYRVIHFIA
 PNOV1: 61 WWKYILLGLADVEANYLIVRAYQYTTLTsvQLLDCFGIPVLMALSWFilyARYRVIHFIA 120

 HSNV1: 168 VAVCLLGVTMVGADILAGREDNSGSDVLIGDILVLLGASLYAISNVCEEYIVKKLSRQE 227
 VAVCLLGVTMVGADILAGREDNSGSDVLIGD+LVLLGASLYA+SNVCEEYIVKKLSRQE
 PNOV1: 121 VAVCLLGVTMVGADILAGREDNSGSDVLIGDVLVLLGASLYAVSNVCEEYIVKKLSRQE 180

 HSNV1: 228 FLGMVGLFGTIISGIQLLIVEYKDIAIHWDWKIALLFVAFALCMFCLYsfMPLVIKvTS 287
 FLGMVGLFGTIISGIQLLIVEYKDIAIHWDWKIALLFVAFALCMFCLYsfMPLVIKvTS
 PNOV1: 181 FLGMVGLFGTIISGIQLLIVEYKDIAIHWDWKIALLFVAFALCMFCLYsfMPLVIKvTS 240

 HSNV1: 288 ATSVNLGILTADLYSLFVGLFLFGYKFSGLYILSFTVIMVGFILYcSTPTrTAEPaESSV 347
 ATSVNLGILTADLYSLF GLFLFGYKFSGLYILSF VIMVGFILYcSTPTrTAEPaESSV
 PNOV1: 241 ATSVNLGILTADLYSLFFGLFLFGYKFSGLYILSFAVIMVGFILYcSTPTrTAEPaESSV 300

 HSNV1: 348 -PPVTSIGIDNLGLKLEENLQETHSAVL 374
 PPVTSIGIDNLGLKLEENL ETHS L
 PNOV1: 301 PPPVTSIGIDNLGLKLEENLPETHSVAL 328

Fig. 32

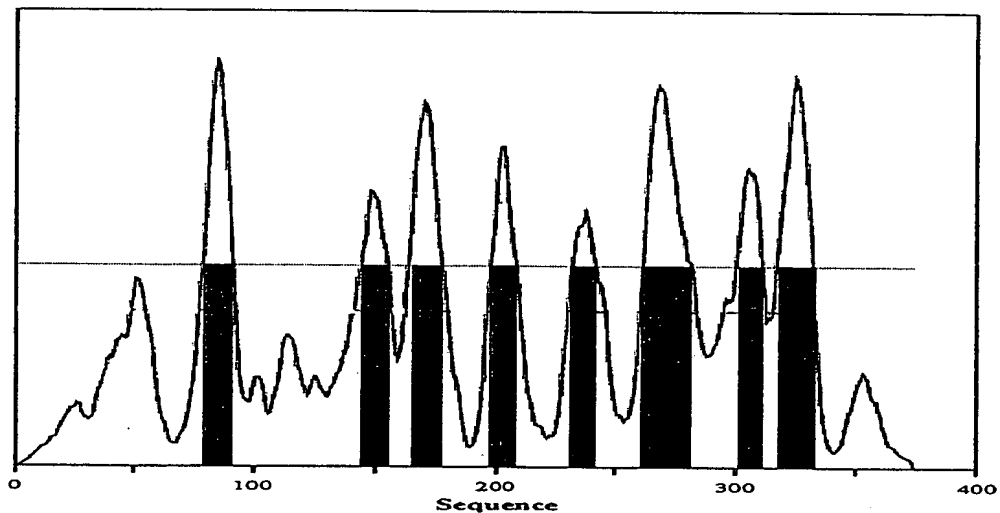


Fig. 33

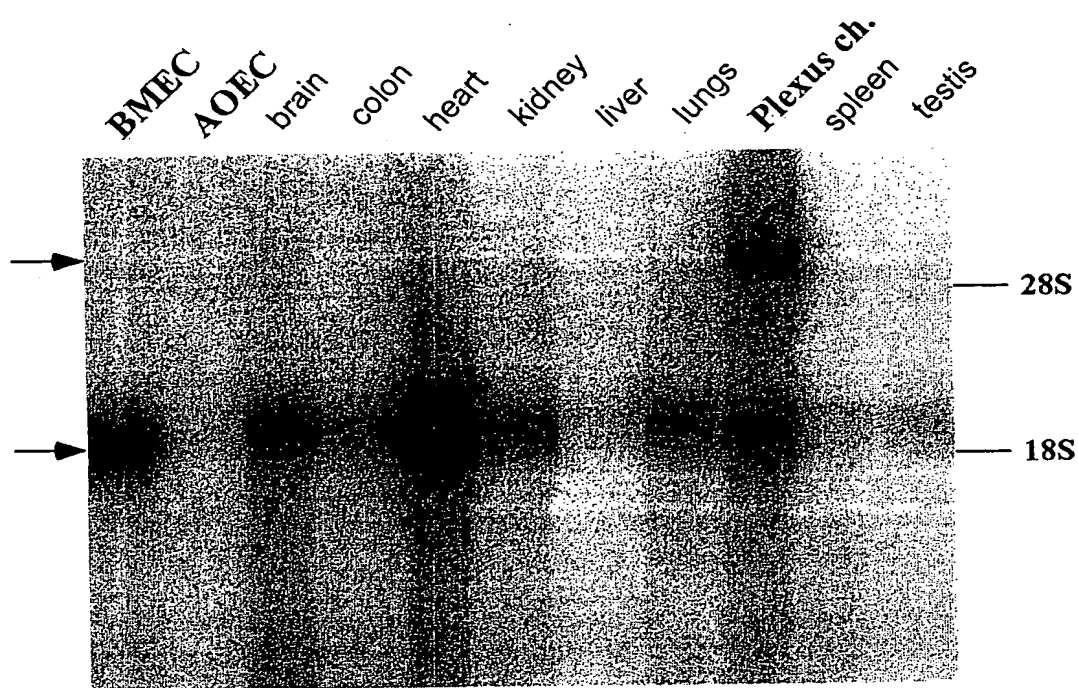


Fig. 34

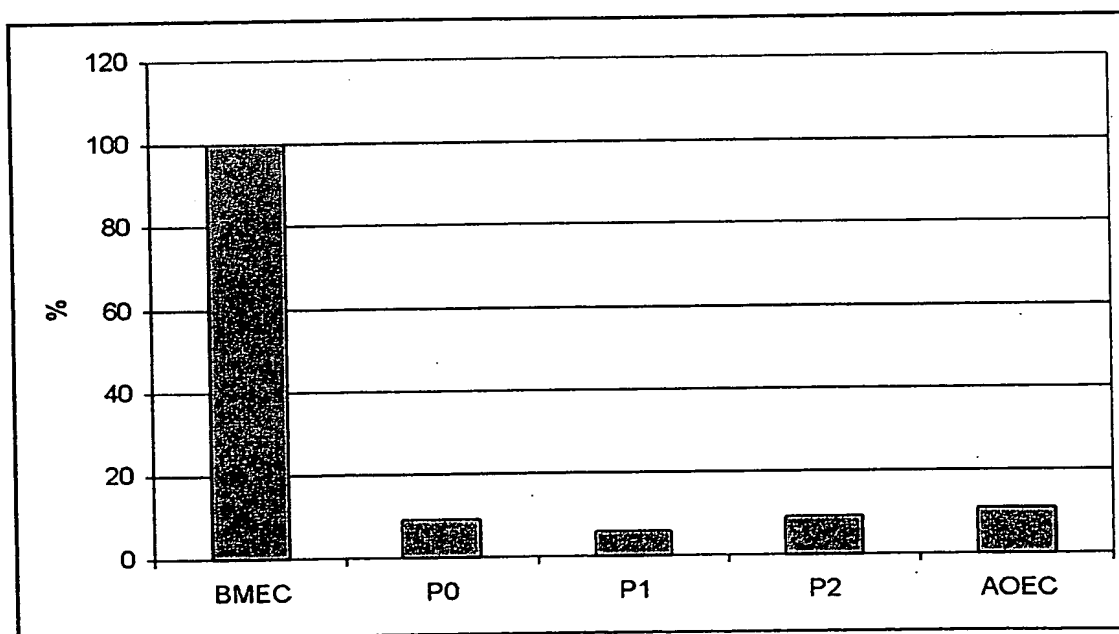


Fig. 35

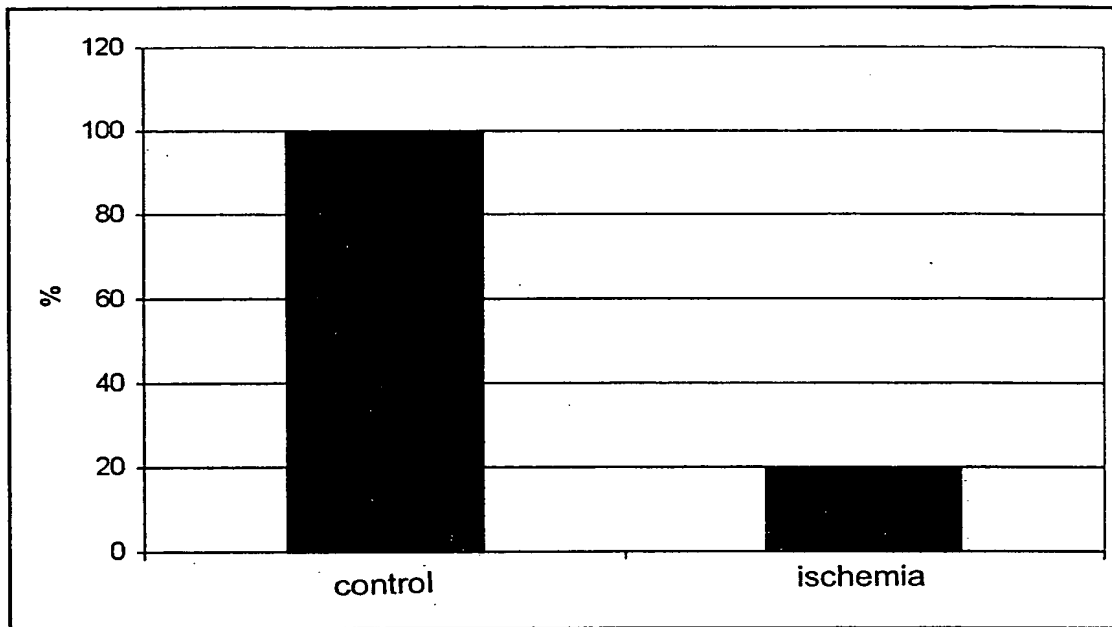


Fig. 36

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